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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/771,536B

DATE: 02/05/2002
 TIME: 11:47:51

Input Set : A:\Sequence Listing.txt
 Output Set: N:\CRF3\02052002\I771536B.raw

ENTERED

3 <110> APPLICANT: Langridge, William H. R.
 4 Yu, Jie
 5 Arakawa, Takeshi
 7 <120> TITLE OF INVENTION: Transgenic Plant-Based Vaccines
 9 <130> FILE REFERENCE: 12273-3
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/771,536B
 12 <141> CURRENT FILING DATE: 2001-01-29
 14 <160> NUMBER OF SEQ ID NOS: 15
 16 <170> SOFTWARE: PatentIn version 3.1
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 376
 20 <212> TYPE: DNA
 21 <213> ORGANISM: Vibrio cholerae
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 28 ctaaatgata agatattgtc gtatacagaa tctctagctg gaaacagaga gatggctatc 180
 30 attactttta agaatgggtgc aacttttcaa gtagaagtac caggtagtca acatatagat 240
 32 tcacaaaaaa aagcgattga aaggatgaag gataccctga ggattgcata tcttactgaa 300
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 41 <212> TYPE: DNA
 42 <213> ORGANISM: artificial sequence
 44 <220> FEATURE:
 45 <223> OTHER INFORMATION: completely synthesized
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 64 <211> LENGTH: 391
 65 <212> TYPE: DNA
 66 <213> ORGANISM: artificial sequence
 68 <220> FEATURE:
 69 <223> OTHER INFORMATION: completely synthesized hinge coding region and Vibrio
 cholerae

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76 ctaaagata agatattgtc gtatacagaa tctctagctg gaaacagaga gatggctatc    180
78 attactttta agaatggtgc aacttttcaa gtagaagtac caggtagtca acatatagat    240
80 tcacaaaaaa aagcgattga aaggatgaag gataccctga ggattgcata tcttactgaa    300
82 gctaaagtcg aaaagttatg tgtatggaat aataaaacgc ctcatgcatg tgccgcaatt    360
84 agtatggcaa attggcccag gcccgggata a                                     391
87 <210> SEQ ID NO: 5
88 <211> LENGTH: 54
89 <212> TYPE: DNA
90 <213> ORGANISM: Vibrio cholerae
92 <400> SEQUENCE: 5
93 atggtaaaga taatatttgt gttttttatt ttcttatcat cattttcata tgca          54
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97 <211> LENGTH: 24
98 <212> TYPE: DNA
99 <213> ORGANISM: artificial sequence
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102 <223> OTHER INFORMATION: completely synthesized
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109 <211> LENGTH: 27
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111 <213> ORGANISM: artificial sequence
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114 <223> OTHER INFORMATION: completely synthesized
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121 <211> LENGTH: 66
122 <212> TYPE: DNA
123 <213> ORGANISM: Rotavirus sp.
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128 ttgact                                                                66
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161 <223> OTHER INFORMATION: Vibrio cholerae and Rotavirus sp.
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166 ggaacacctc aaaatattac tgatttgtgt gcagaatacc acaacacaca aatacatatc      120
168 cttaatgata agatattgtc gtatacagaa tctctagctg gaaacagaga gatggctatc      180
170 attactttta agaatggtgc aacttttcaa gtagaagtac caggtagtca acatatagat      240
172 tcacaaaaaa aagcgattga aaggatgaag gataccctga ggattgcata tcttactgaa      300
174 gctaaagtcg aaaagttatg tgtatggaat aataaaacgc ctcatgcgat tgccgcaatt      360
176 agtatggcaa attggccag gcccgggaga gctcgataag ttgactacta gggagattga      420
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184 <211> LENGTH: 444
185 <212> TYPE: DNA
186 <213> ORGANISM: Escherichia coli
188 <400> SEQUENCE: 12
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193 gaaagttaca gagtaatgac tcaagttcat acaaacgatg caactaaaaa agtaattggt      180
195 aaacttgctg atacaccaca gcttacagat gttctgaatt caactgttca aatgcctatc      240
197 agtgtgtcat ggggaggaca agtattatct tctacaacag ccaaagaatt tgaagctgct      300
199 gctttgggat attctgcac cgggtgaaat ggcgtatcat cttctcaaga gttagtaatt      360
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208 <212> TYPE: DNA
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214 caatctaaag ttaaagaca aatattttca ggctatcaat ctgatattga tacacataat      120
216 agaattaaag atgagttgtg a                                         141
219 <210> SEQ ID NO: 14
220 <211> LENGTH: 54
221 <212> TYPE: DNA
222 <213> ORGANISM: Vibrio cholerae
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230 <212> TYPE: DNA

231 <213> ORGANISM: artificial sequence

233 <220> FEATURE:

234 <223> OTHER INFORMATION: Vibrio cholerae and Escherichia coli

236 <400> SEQUENCE: 15

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237 atggtaaaga taatatttgt gtttttttatt ttcttatcat cattttcata tgcagtcgac      60
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241 gatggcaatg ctctgccatc agctgtaaag ttagcttatt ctcccgcac aaaaactttt      180
243 gaaagttaca gagtaatgac tcaagttcat acaaacgatg caactaaaaa agtaattggt      240
245 aaacttgctg atacaccaca gcttacagat gttctgaatt caactgttcaaatgcctatc      300
247 agtgtgtcat ggggaggaca agtattatct tctacaacag ccaaagaatt tgaagctgct      360
249 gctttgggat attctgcac cggtgtaaag ggcgtatcat cttctcaaga gttagtaatt      420
251 agcgtgcac ctaaaactgc cggtaccgcc ccaactgcag gaaactattc aggagtagta      480
253 tctcttgtaa tgactttggg atccgtcgac atcagtaata cttgcgatga aaaaacccaa      540
255 agtctaggtg taaaattcct tgacgaatac caatctaaag ttaaagaca aatattttca      600
257 ggctatcaat ctgatattga tacacataat agaattaaag atgagttgtg a              651
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VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number